

TGTCTGCGCTGCGATGAG

5'	GCGCC	ATC	AGC	CGC	CGC	GGA	GAT	ATC	CGC	CGG	GGG	AGA	ATA	GGG	TTG	CAC	CAT	CCC
	11		20		29					38				47				56
	65		74		83					92				101				110
	119		128		137					146				155				164
TGT	CCG	CCC	TTC	CCC	TAG	CGT	TAC	TTC	CGG	TCC	CTC	GCT	GAG	GGG	GTT	CGT	CGC	
	173		182		191					200				209				218
GCT	CCC	AGG	AGG	CGT	GAA	CGG	ACC	ATG	AGC	GTG	GGC	TTC	ATC	GGG	GCC	GCC		
		M	S	V	G	F	I	G	A	G								
	227		236		245					254				263				272
CAG	CTG	GCC	TAT	GCT	CTG	GCG	CGG	GGC	TTC	ACG	GCA	GGC	ATC	CTG	TCG	GCT		
Q	L	A	Y	A	L	A	R	G	F	T	A	A	G	I	L	S	A	
	281		290		299					308				317				326
CAC	AAG	ATA	ATA	GCC	AGC	TCC	CCA	GAA	ATG	AAC	CTG	CCC	ACG	GTG	TCC	GCG	CTC	
H	K	I	I	A	S	S	P	E	M	N	L	P	T	V	S	A	L	
	335		344		353					362				371				380
AGG	AAG	ATG	GGT	GTG	AAC	CTG	ACA	CGC	AGC	AAC	AAG	GAG	ACG	GTG	AAG	CAC	AGC	
R	K	M	G	V	N	L	T	S	N	K	E	T	V	K	H	S		
	389		398		407					416				425				434
GAC	GTC	CTG	TTT	CTG	GCT	GTG	AAG	CCA	CAT	ATC	ATC	CCC	TTC	ATC	CTG	GAT	GAG	
D	V	L	F	L	A	V	K	P	H	I	I	P	F	I	L	D	E	

FIGURE 1A

TUNESLOUZI ET AL.

443	ATT	GGG	GCC	GAC	GTG	CAA	GCC	AGA	CAC	ATC	GTG	GTC	TCC	TGT	GCG	GCT	GGT	GTC
I	G	A	D	V	Q	A	R	H	I	V	V	S	C	A	A	G	V	
497	ACC	ATC	AGC	TCT	GTG	GAG	AAG	CTG	ATG	GCA	TTC	CAG	CCA	GCC	CCC	AAA	GTC	
T	I	S	S	V	E	K	R	L	M	A	F	Q	P	A	P	K	V	
551	ATT	CGC	TGC	ATG	ACC	AAC	ACA	CCT	GTG	GTA	GTG	CAG	GAA	GGC	GCT	ACA	GTG	TAC
I	R	C	M	T	N	T	P	V	V	V	Q	E	G	A	T	V	Y	
605	GCC	ACG	GGC	ACC	CAT	GCC	CTG	GTG	GAG	GAT	GGG	CAG	CTC	CTG	GAG	CAG	CTC	ATG
A	T	G	T	H	A	L	V	E	D	G	Q	L	L	E	Q	L	M	
659	AGC	AGC	GTG	GGC	TTC	TGC	ACT	GAG	GTG	GAA	GAG	GAC	CTC	ATC	GAT	GCC	GTC	ACG
S	S	V	G	F	C	T	E	V	E	D	L	I	D	A	V	T		
713	GGG	CTC	AGT	GGC	AGC	GGG	CCT	GCC	TAT	GCA	TTC	ATG	GCT	CTG	GAC	GCA	TTG	GCT
G	L	S	G	S	G	P	A	Y	A	F	M	A	L	D	A	L	A	
767	GAT	GGT	GGG	GTG	RAG	ATG	GGT	TTG	CCA	CGG	CGC	CTG	GCA	ATC	CAA	CTC	GGG	GCC
D	G	G	V	K	M	G	L	P	R	R	L	A	I	Q	L	G	A	

FIGURE 1B

የጥቃት ማረጋገጫ እና ተግባራዊነት

Q	A	L	G	A	A	K	M	L	L	D	S	E	Q	H	P	C
Q	L	K	D	N	V	C	S	P	G	G	A	T	I	H	A	L
F	L	E	S	G	G	F	R	S	L	L	I	N	A	V	E	A
C	I	R	T	R	E	L	Q	S	M	A	D	Q	E	K	I	S
S	T	L	T	P	S	S	P	G	K	L	T	R	S	L	A	V
G	G	K	K	D												
821																
830																
839																
CAG GCT TTG CTG GGA GCT GCC AAG ATG CTG CTG GAC TCG GAG CAG CAT CCA TGC																
925																
884																
893																
CAG CTT AAG GAC AAT GTC TGC TCC CCT GGG GGA GCC ACC ATC CAC GCC CTG CAC																
929																
938																
947																
TTT CTA GAG AGT GGG GGC TTC CGC TCT CTG CTC ATC ATT GCA GTT GAG GCC TCC																
983																
992																
TGT ATC CGA ACA CGA GAG CTA CAG TCC ATG GCC GAC CAA GAA AAG ATC TCC CCA																
1037																
1046																
1055																
GCT GCC CTT AAG AAG ACC CTC TTA GAC AGA GTG AAG CTG GAA TCC CCC ACA GTC																
1091																
1100																
1109																
TCC ACA CTG ACC CCC TCC AGC CCA GGG AAG CTC CTC ACA AGA AGC CTG GCC CTG																
1145																
1154																
GGA GGC AAG AAG GAC TAA GGC AGC ATC TGT CCC CTC TGT GAT TCA GAG CCC TTA																

FIGURE 1C

TRANSCRIPTION

1199	1208	1217	1226	1235	1244
GTT GAG AGC CCC TGC CGC CCC TGC CAC CCC CCT GCC CCG CTC CCA CCA TTG CCC					
1253	1262	1271	1280	1289	1298
CTC CTC AGC TGT GCA AGG AGA AAG CAT GCT TAG GAA GTT TTC AGG TCC TTG TGA					
1307	1316	1325	1334	1343	1352
TAA AAC CTC CTT AAA TCT GTT CAG ACC AAG CAA TGC GAG CCT CCT CTC CTG TCC					
1361	1370	1379	1388	1397	1406
CAT GTT GGA AGT TGC TCT GAA GGG GTG GTA GAT GCT GGA AGC CAG ACA CAA CCC					
1415	1424	1433	1442	1451	1460
TGC GTA CGC TGC TCA GTT GGT GGA GAC TGG GGC TGG GAC TGG AGT CAG CCC AGC					
1469	1478	1487	1496	1505	1514
TGG GAG GAG GGG CTG GGG AGG ATC TGC AGC TGA AGC CCG AGG CAG GGT TGG TGT					
1523	1532	1541	1550	1559	1568
GAT GCC AAG GCA AAG TGG TGA GGA GAA AAC AGG AAA CGG GCT TTC TCT GAA TTG					
1577	1586	1595	1604	1613	1622
GTA AAT GGG AAA GAA GTG AGC AAC TTA AGA TTG TCA CAA TTA ATC ACA AGT GTA					
1631	1640	1649	1658	1667	1676
CAG GAT TAG ACT GGG TTT ATA TTT AAC TCT TGC TTC ATA GGT GTA CCA TTT AAA					

FIGURE 1D

FIGURE 1E

T G T C A G C T T T G G G

1685 1694 1703 1712 1721 1730
GAG TGT TAT TTA ATG CTA AGT TTA ACT GCT TTA ATA AAG TTT ATT TTT AAA TAT
1739
CAA AAA AAA AAA 3'

1	M S V G F I G A G Q L A Y A L A R G F T A A G I L S A H K I I A S S P	2278458
1	M S V G F I G A G Q L A F A L A K G F T A A G V L A A H K I M A S S P	g189498
36	E M N L P T V S A L R K M G V N L T R S N K E T V K H S D V L F L A V	2278458
36	D M D L A T V S A L R K M G V K L T P H N K E T V Q H S D V L F L A V	g189498
71	K P H I I P F I L D E I G A D V Q A R H I V V S C A A G V T I S S V E	2278458
71	K P H I I P F I L D E I G A D I E D R H I V V S C A A G V T I S S S T E	g189498
106	K L M A F Q P A P K V I R C M T N T P V V V Q E G A T V Y A T G T H	2278458
106	K K L S A F R P A P R V I R C M T N T P V V V R E G A T V Y A T G T H	g189498
141	A L V E D G Q L L E Q L M S S V G F C T E V E E D L I D A V T G L S G	2278458
141	A Q V E D G R L M E Q L L S T V G F C T E V E E D L I D A V T G L S G	g189498
176	S G P A Y A F M A L D A L A D G G V K M G L P R R L A I Q L G A Q A L	2278458
176	S G P A Y A F T A L D A L A D G G V K M G L P R R L A V R L G A Q A L	g189498
211	L G A A K M L L D S E Q H P C Q L K D N V C S P G G A T I H A L H F L	2278458
211	L G A A K M L L H S E Q H P G Q L K D N V S S P G G A T I H A L H V L	g189498

FIGURE 2A

246	E S G G F R S L L I N A V E A S C I R T R E L Q S M A D Q E K I S P A	2278458
246	E S G G F R S L L I N A V E A S C I R T R E L Q S M A D Q E Q V S P A	g189498
281	A L K K T L L D R V K L E S P T V S T L T P S S P G K L L T R S L A L	2278458
281	A [T] K K T [I] L D [K] V K L D S P A G T A L S P S G H T K L L P R S L A P	g189498
316	G G K K D	2278458
316	A [G K [-D]	g189498

FIGURE 2B